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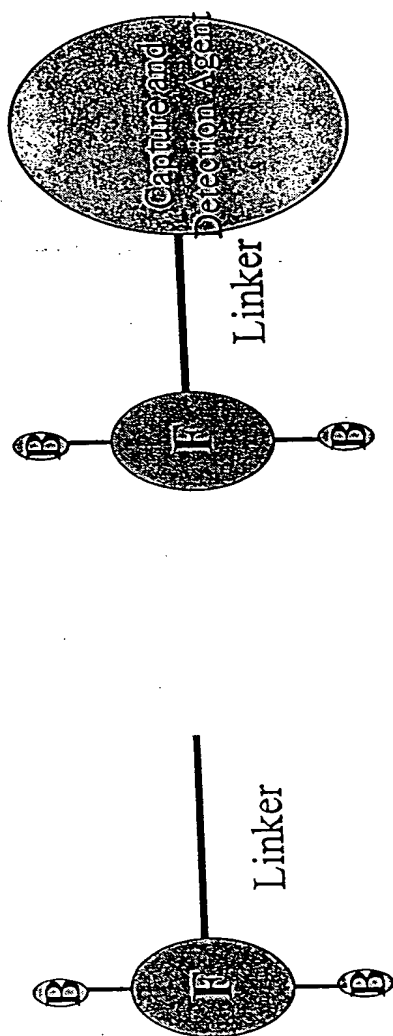
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High-throughput Target ID



Library of Target ID Compounds

Library of Bioactive Compounds

Use corresponding activity-based probe to identify the biological target

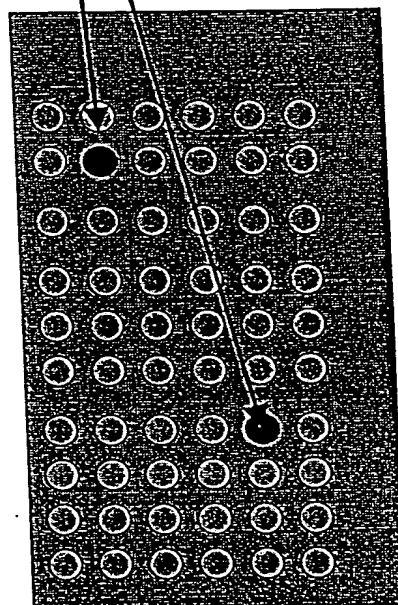
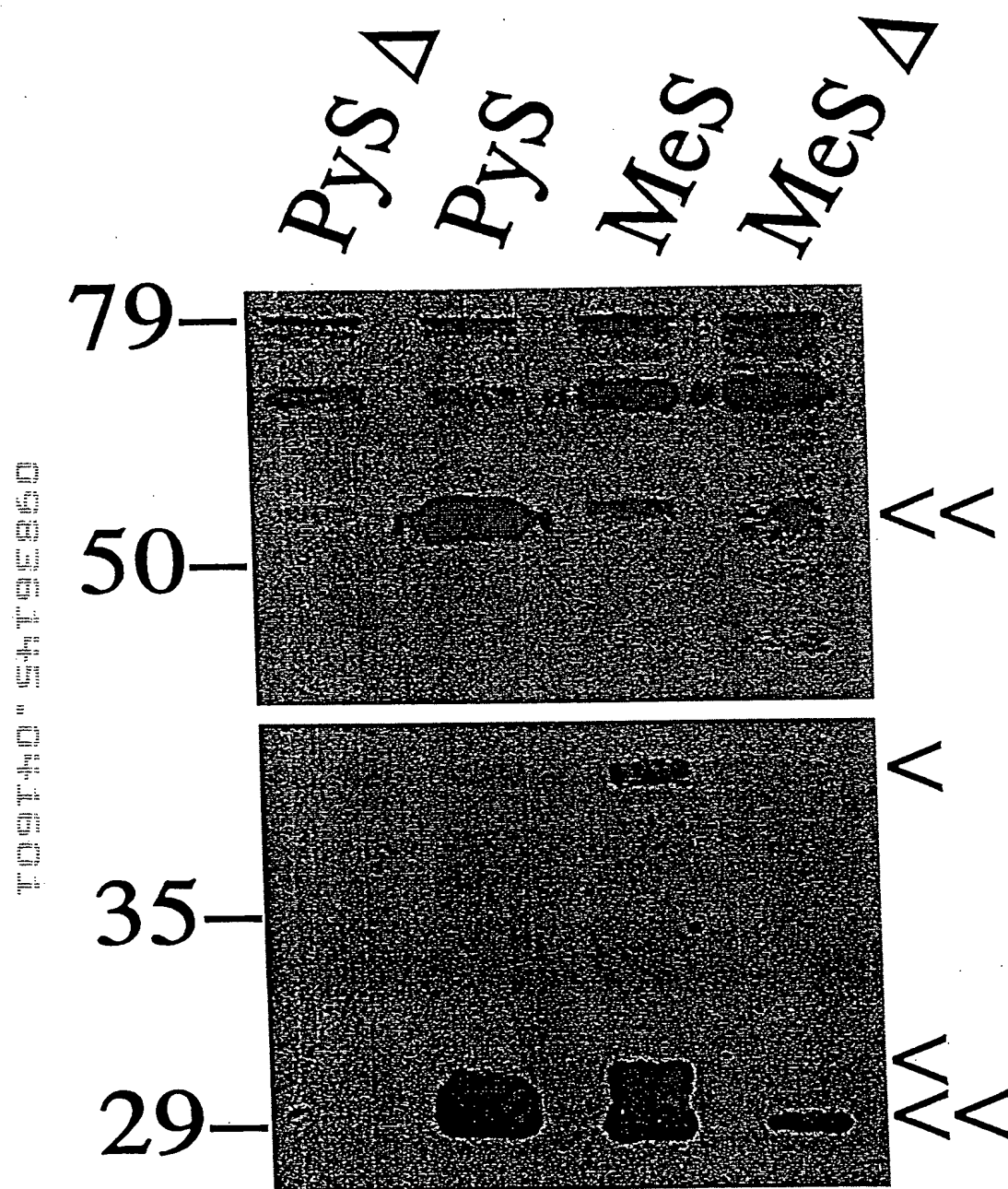


FIGURE 1



Non-Directed Tagged Library of Sulfonates Identifies Probe for ADH Superfamily of Enzymes

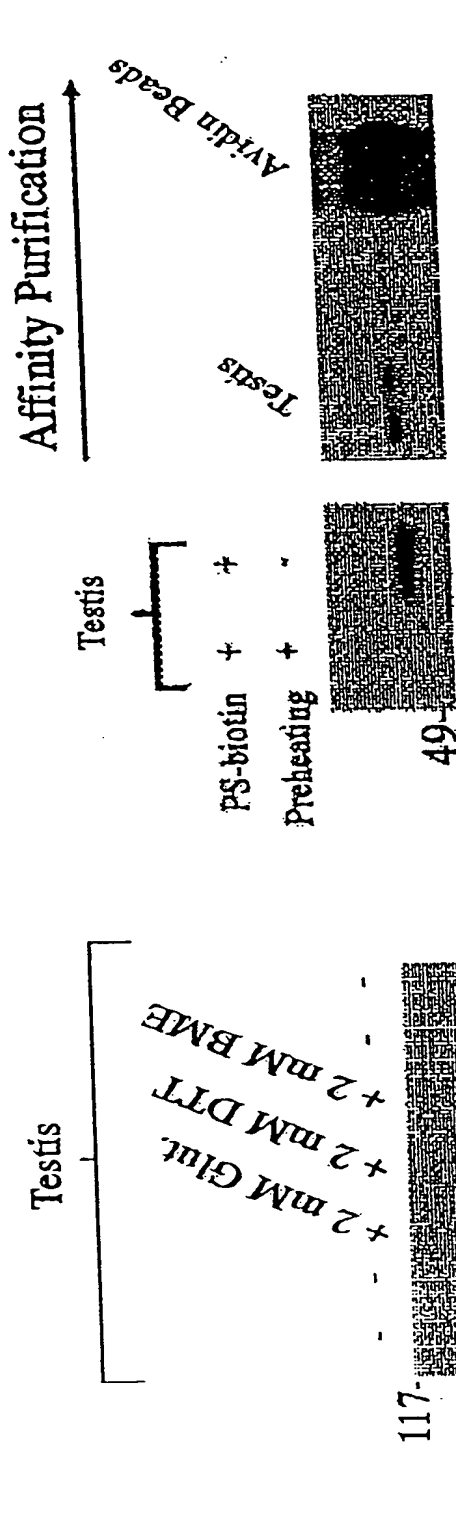
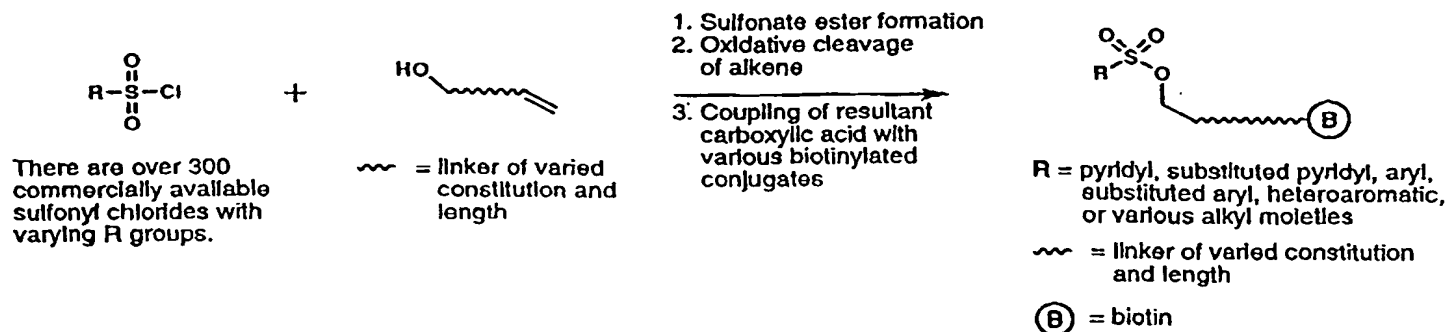


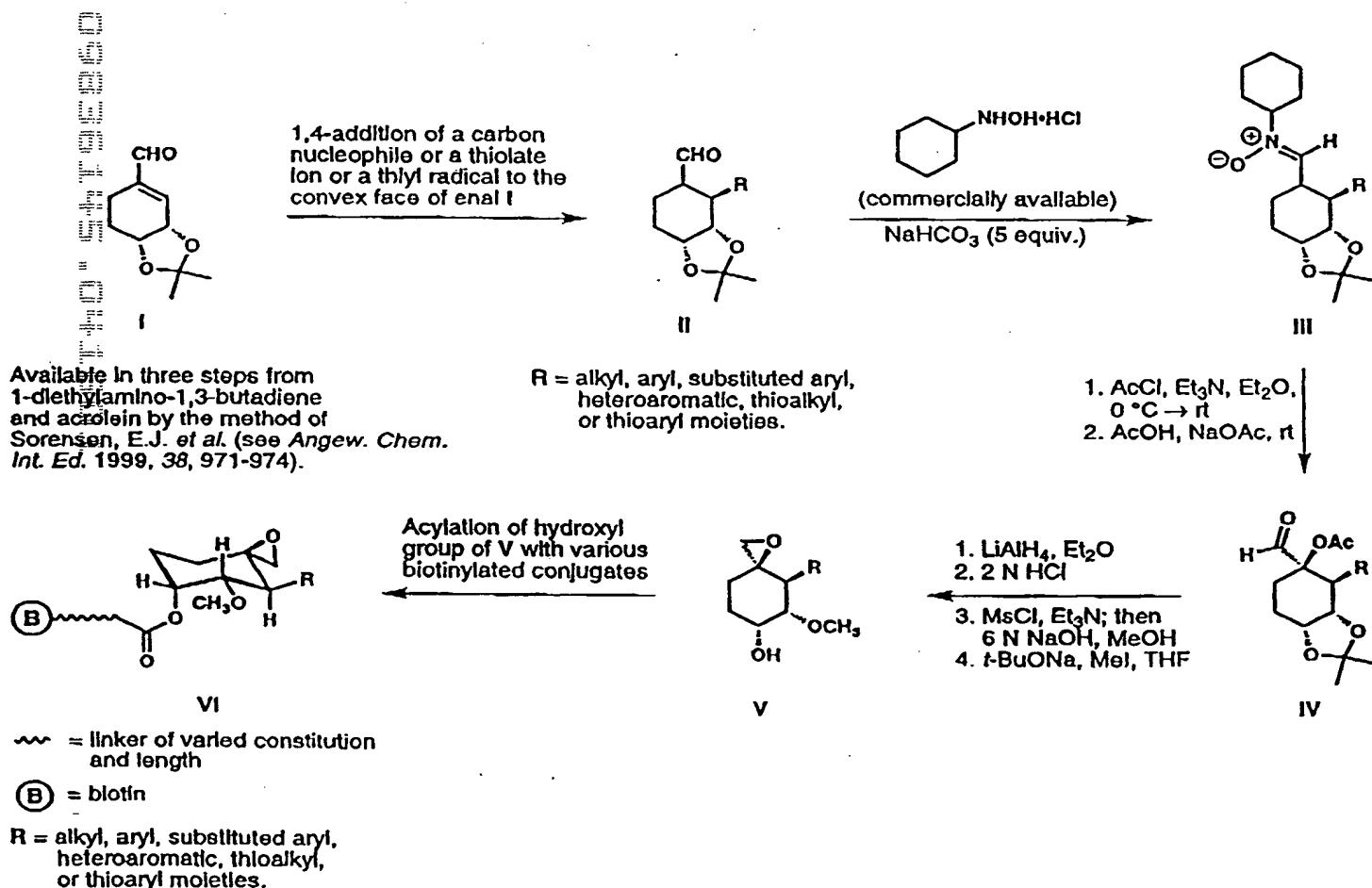
FIGURE 3

- MALDI mapping identifies tagged protein as aldehyde dehydrogenase (ADH, cytosolic class II)
- 28 ADHs in fly genome
 - Involved in retinoic acid biosynthesis and catabolism of alcohol and chemotherapeutic agents

FIGURE 4



Scheme 1. A pathway for syntheses of various biotinylated sulfonate esters for use in activity-based proteomics studies.



Scheme 2. A strategy for convergent, stereocontrolled syntheses of conformationally well-defined spiroepoxides of type VI. Literature precedent for I → II → III → IV → V can be found in Sorensen, E.J. *et al.* *Angew. Chem. Int. Ed.* 1999, 38, 971-974. Compounds of type VI are analogs of the metalloprotease (MetAp-2) inhibitor fumagillin and will be employed as covalent affinity agents in activity-based proteomics studies.

FP-Biotin: a kinetic reporter of SH Activity

The rates at which the majority of SHs react with FP-biotin can be experimentally followed

FP-biotin readily detects low femtomole quantities of SHs directly in complex cell/tissue proteomes

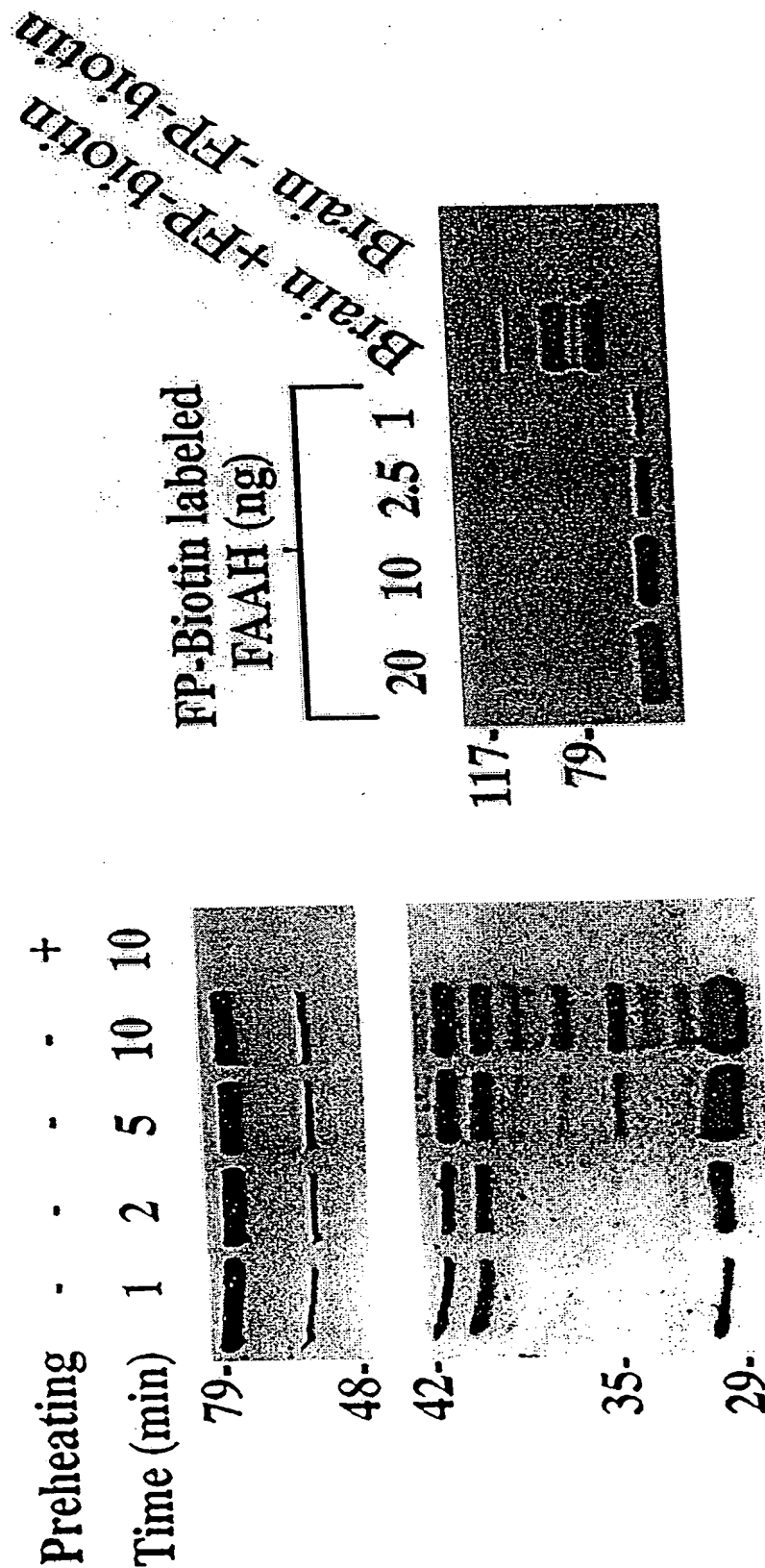


FIGURE 5

Utility of Multiplexed probes in identifying Serine Hydrolases

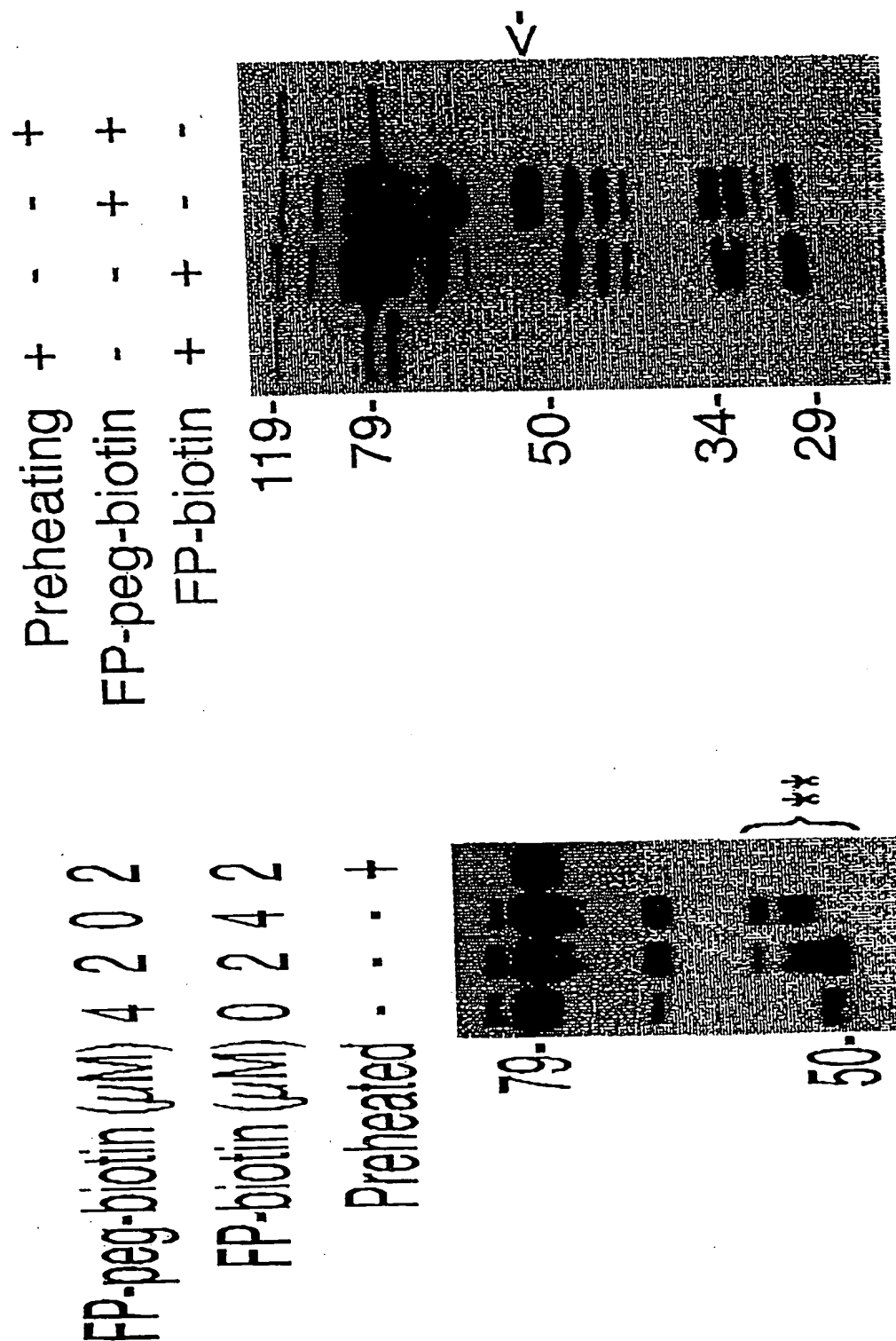


FIGURE 6

FIGURE 7

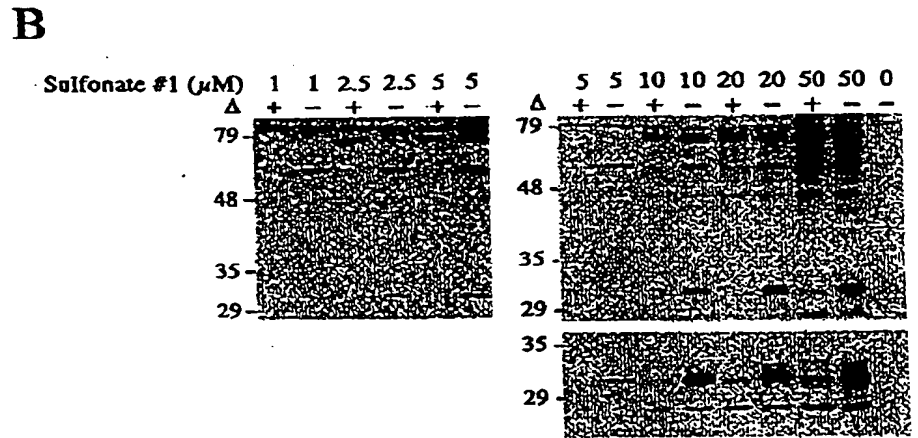
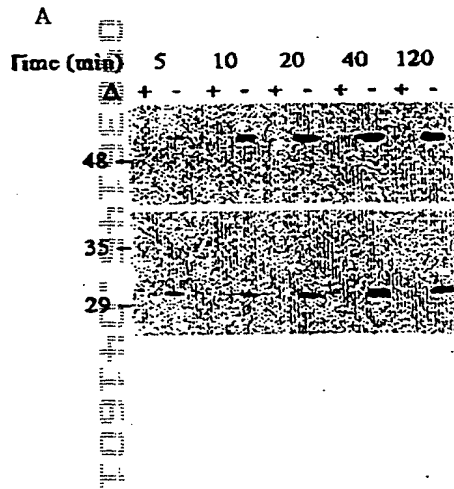
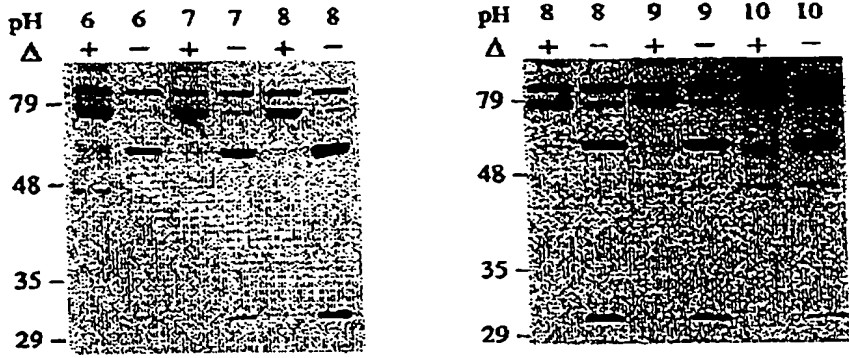
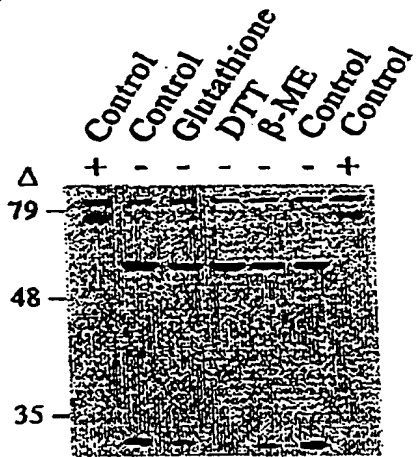


FIGURE 7

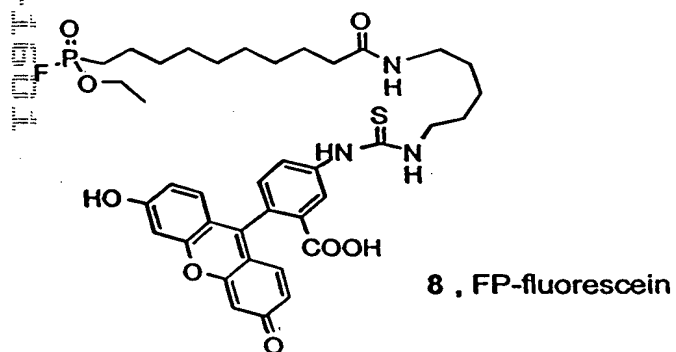
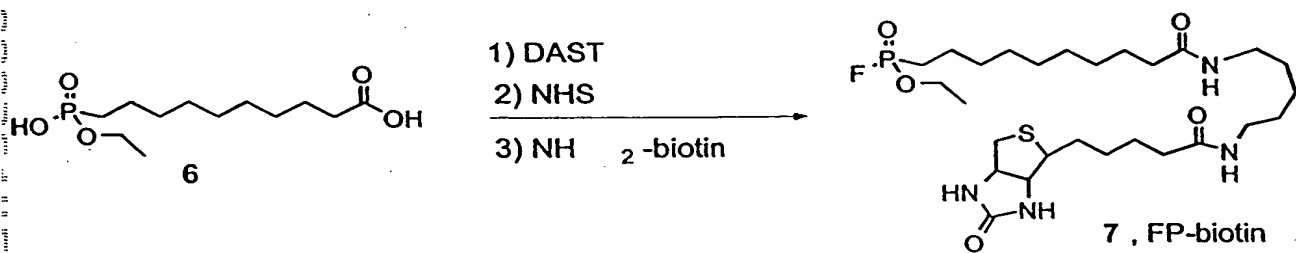
C



D



TsCl $\left\{ \begin{array}{l} \rightarrow 1, X = OH \\ \rightarrow 2, X = OTs \end{array} \right.$
NaI $\left\{ \begin{array}{l} \rightarrow 3, X = I \end{array} \right.$



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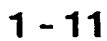
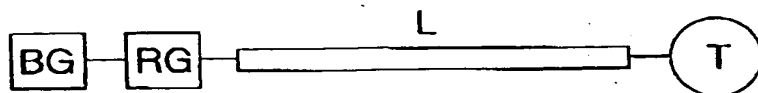
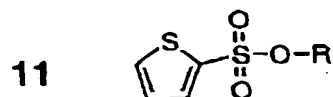
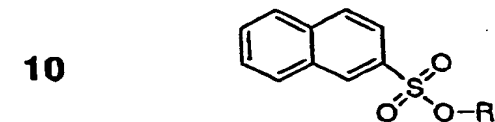
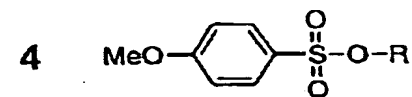
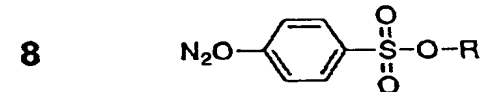
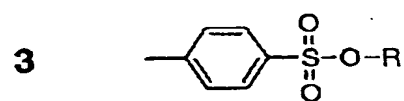
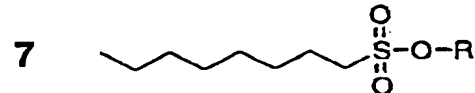
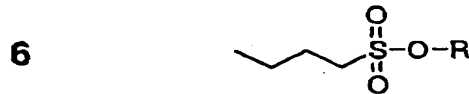
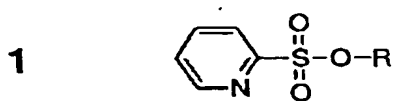
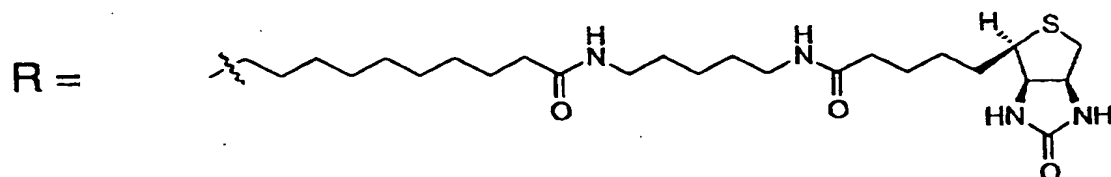


FIGURE 10

A.



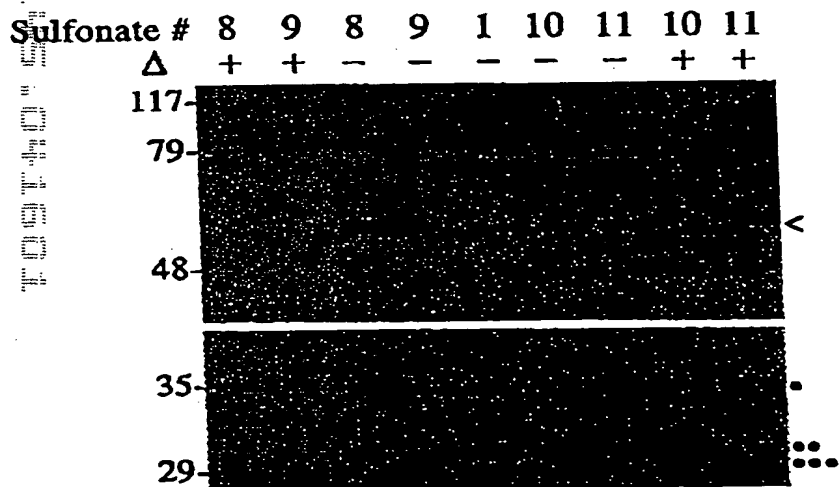
B.



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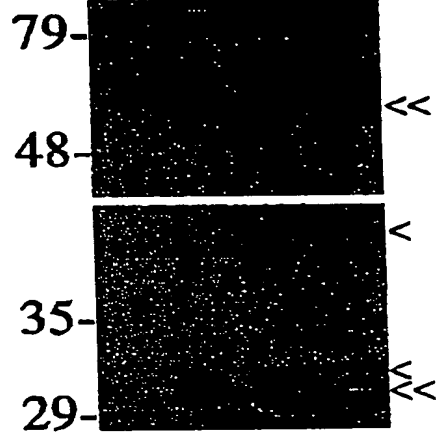
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B

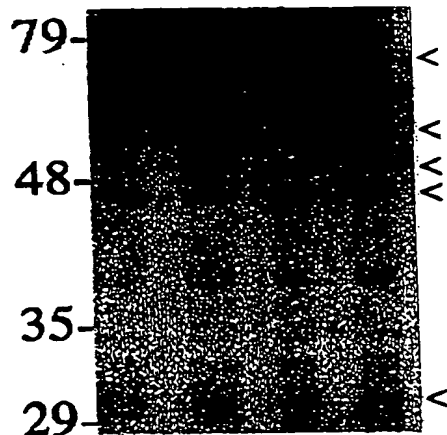


A

Sulfonate # 1 1 5 5

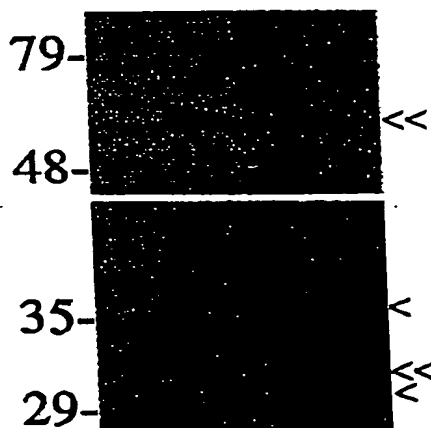
$$\Delta + - - +$$


1 1 5 5

$$+ \quad - \quad - \quad +$$


B

Sulfonate # 1 1 9 9

$$\Delta \quad + \quad - \quad - \quad +$$


1 1 9 9

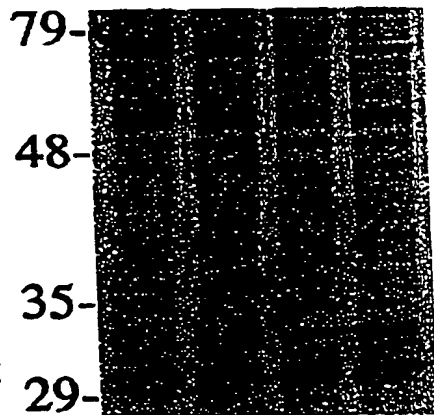
$$+ \quad - \quad - \quad +$$


FIGURE 13

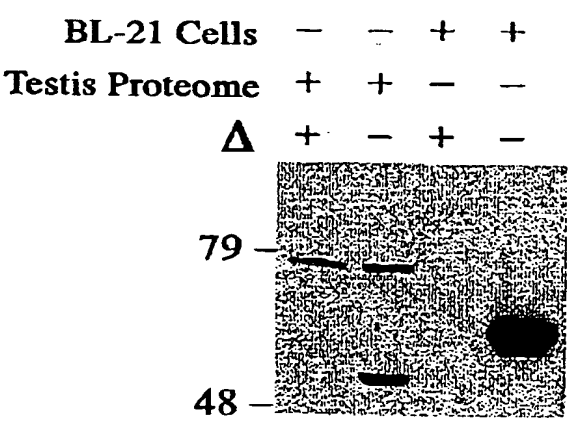
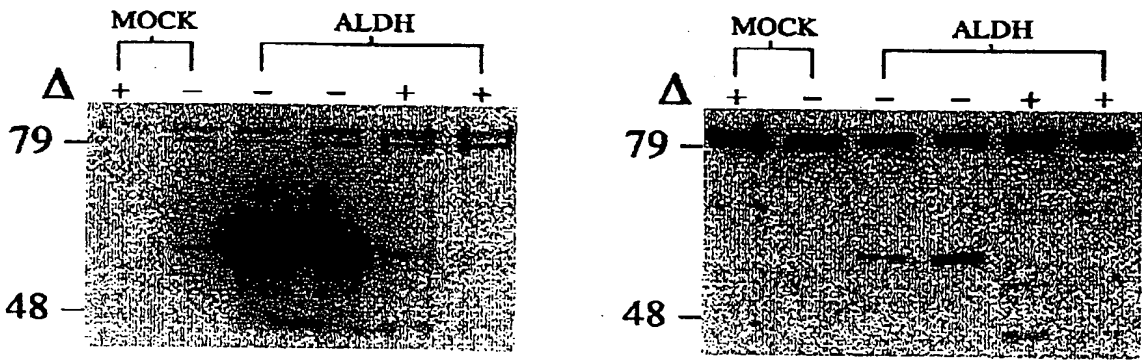
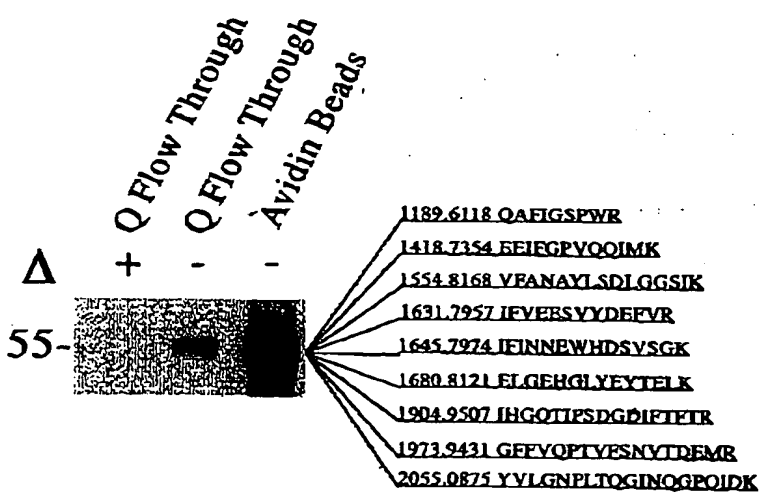
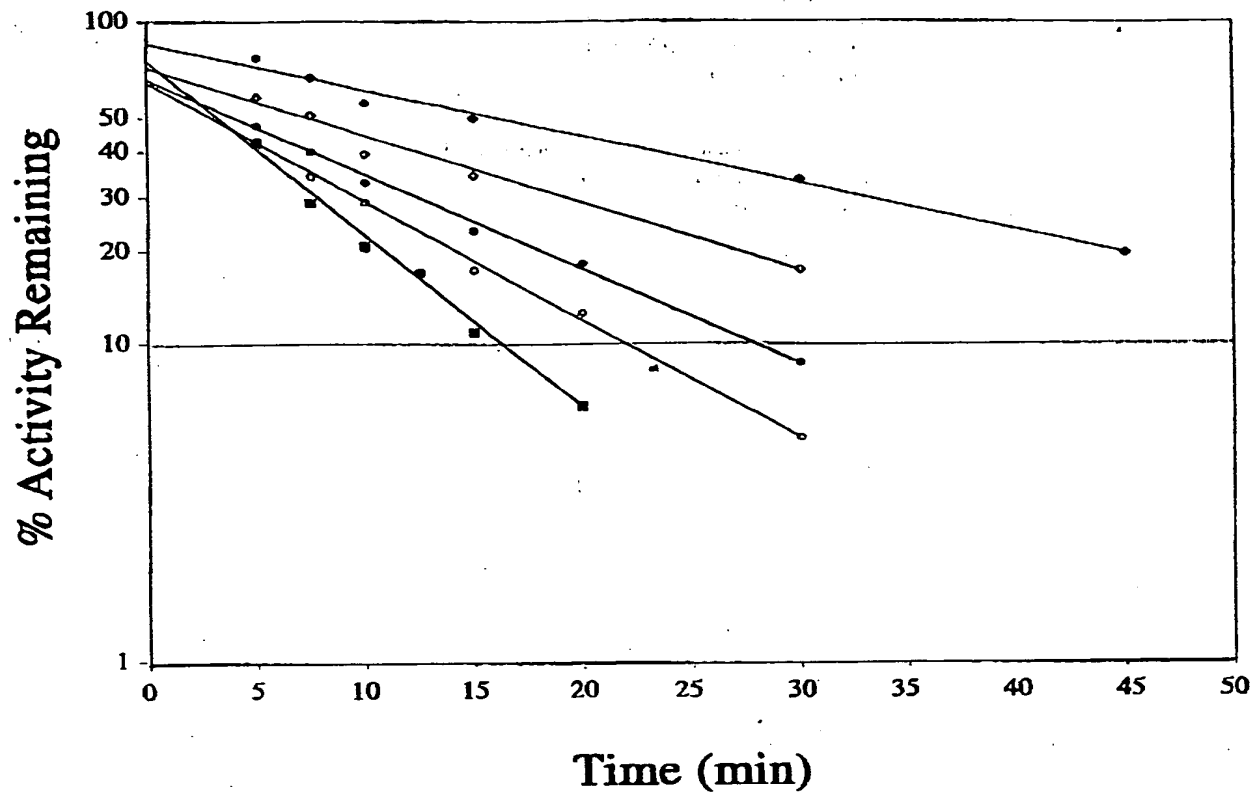


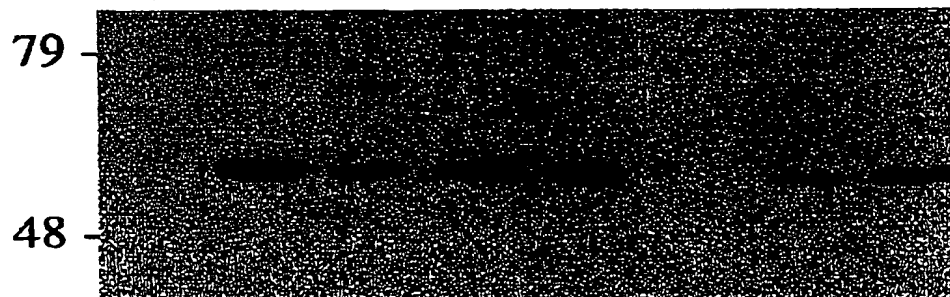
FIGURE 14

A



B

Competitor #	-	-	15	17	16	15	17	16
[Competitor (μ M)]	0	0	5	5	5	50	50	50
Δ	+	-	-	-	-	-	-	-



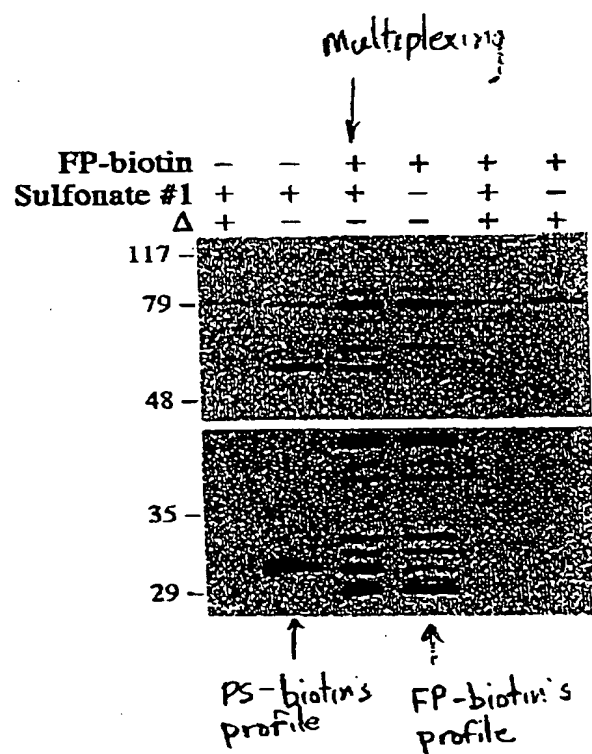
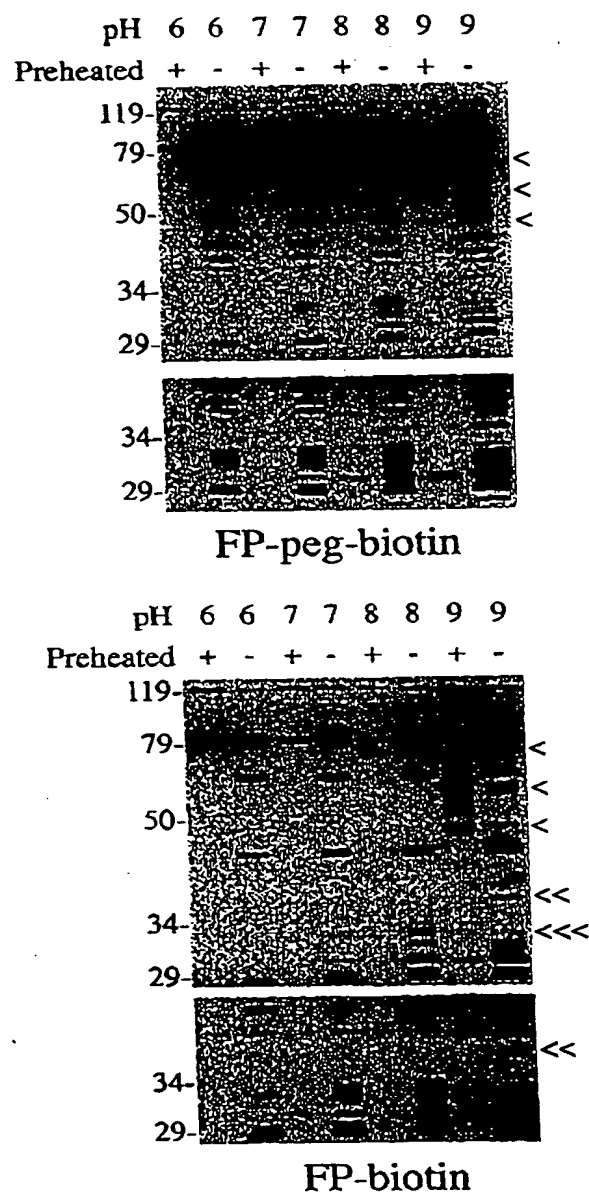
[illegible]

FIGURE 16



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FIGURE 17

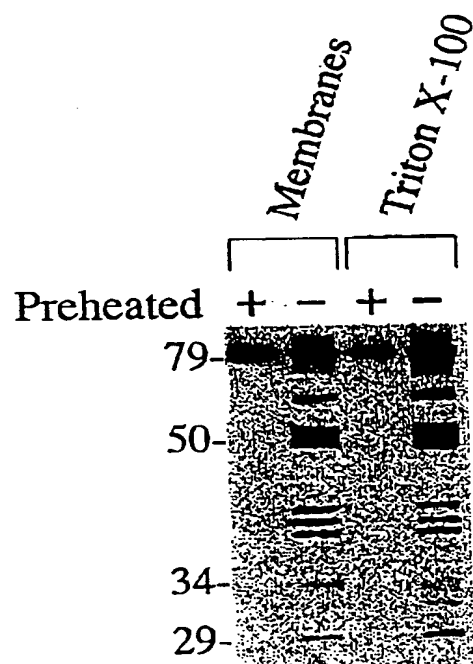
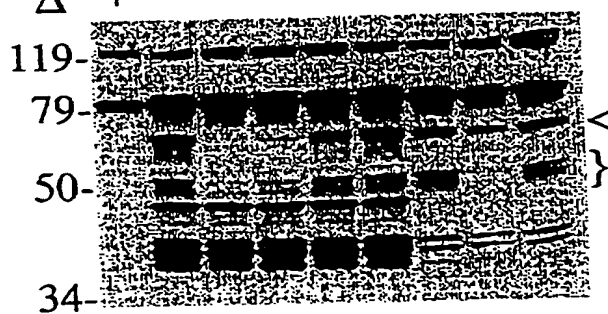


FIGURE 18

FP-peg-biotin	+	+	+	+	+	+	-	-	-
FP-biotin	-	-	-	-	-	-	+	+	+
OTFMK	0	0	200	50	5	1	0	200	50
Δ	+	-	-	-	-	-	-	-	-



0936145-041694

FIGURE 19

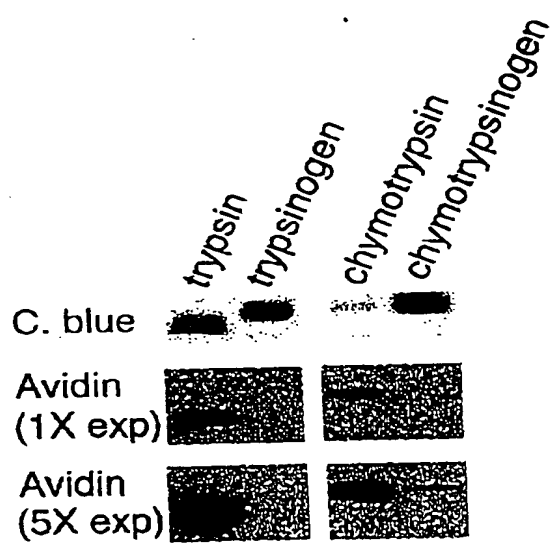
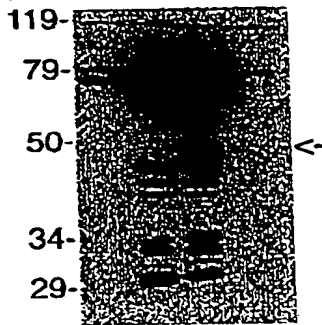


FIGURE 20

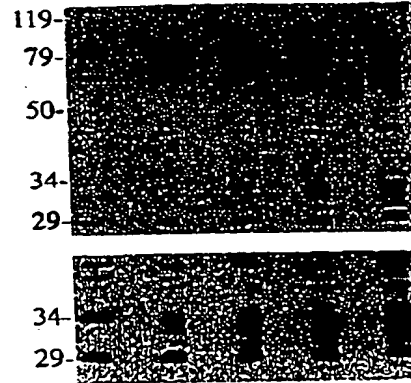
A

FP-peg-biotin	-	-	+	+
FP-biotin	+	+	-	-
Preheated	+	-	-	+



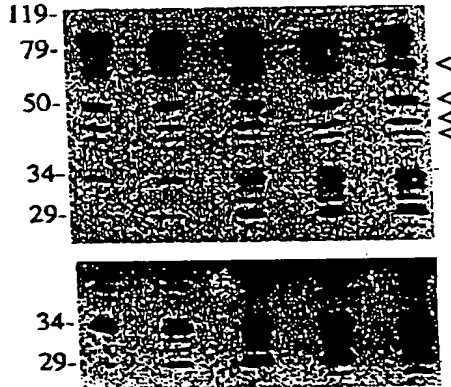
B

FP-biotin (μ M)	0.5	1	1	2	2	4	4	8	8
Preheated	-	+	-	+	-	+	-	+	-



C

FP-peg-biotin (μ M)	0.5	1	1	2	2	4	4	8	8
Preheated	-	+	-	+	-	+	-	+	-



D

FP-peg-biotin (μ M)	1	2	8
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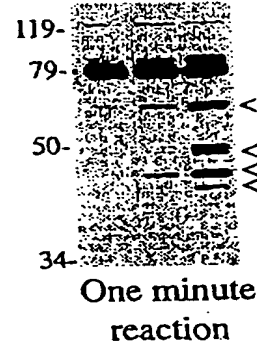


FIGURE 21

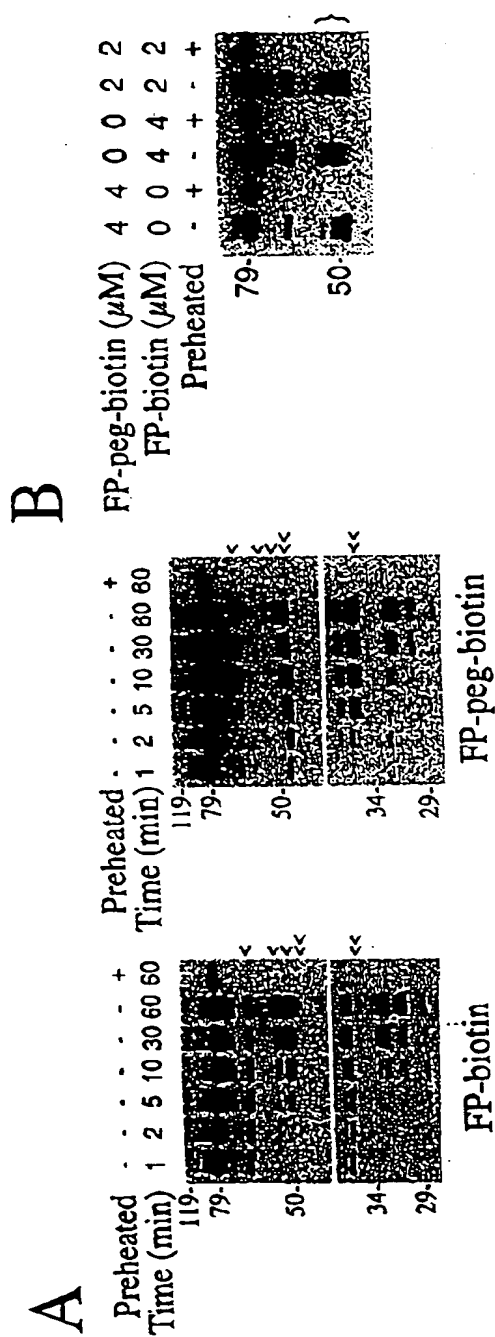


FIGURE 22

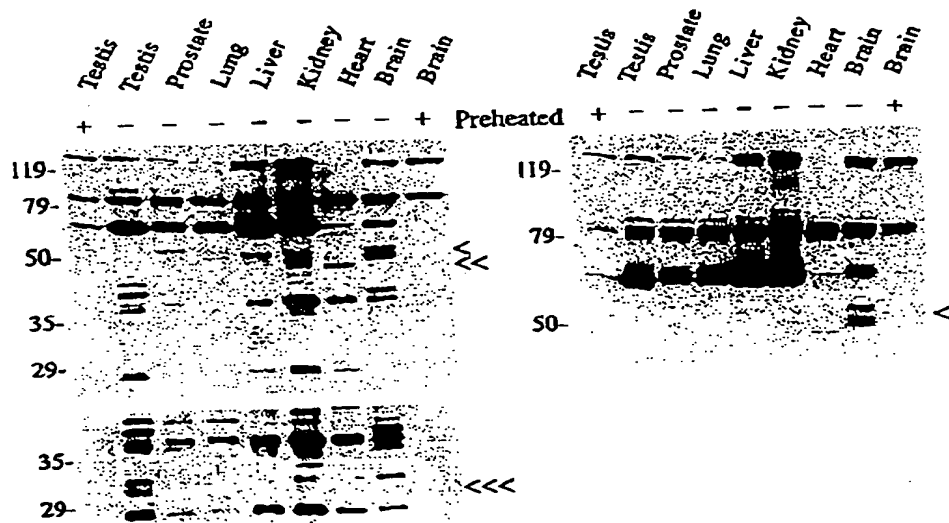


FIGURE 23

